

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

<p>In re Application of:</p> <p>Ashkenazi et al.</p> <p>Serial No.: Not yet assigned</p> <p>Filed: Herewith</p> <p>For: <i>Secreted and Transmembrane Polypeptides and Nucleic Acids Encoding the Same</i></p>	<p>Group Art Unit: Not yet assigned</p> <p>Examiner: Not yet assigned</p>
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**PRELIMINARY AMENDMENT**

Assistant Commissioner of Patents  
Washington, D.C. 20231

Sir:

Prior to substantive examination of the above captioned patent application (which is filed herewith), and for calculation of the proper filing fee, Applicants respectfully request that the following amendments be entered.

**Serial No.:** Not yet assigned

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In the claims:

Please cancel Claims 1-57 without prejudice or disclaimer.

Please add new Claims 58-70 as follows.

–58. (New) An isolated polypeptide having at least 80% amino acid sequence identity to:

(a) the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523);

(b) the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523),

lacking its associated signal peptide;

(c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523);

(d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523), lacking its associated signal peptide; or

(e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209487.

59. (New) The isolated polypeptide of Claim 58 having at least 85% amino acid sequence identity to:

(a) the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523);

(b) the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523),

lacking its associated signal peptide;

(c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523);

(d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523), lacking its associated signal peptide; or

(e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209487.

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60. (New) The isolated polypeptide of Claim 58 having at least 90% amino acid sequence identity to:

(a) the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523);

(b) the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523), lacking its associated signal peptide;

(c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523);

(d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523), lacking its associated signal peptide; or

(e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209487.

61. (New) The isolated polypeptide of Claim 58 having at least 95% amino acid sequence identity to:

(a) the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523);

(b) the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523), lacking its associated signal peptide;

(c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523);

(d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523), lacking its associated signal peptide; or

(e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209487.

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62. (New) The isolated polypeptide of Claim 58 having at least 99% amino acid sequence identity to:

(a) the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523);

(b) the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523), lacking its associated signal peptide;

(c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523);

(d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523), lacking its associated signal peptide; or

(e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209487.

63. (New) An isolated polypeptide comprising:

(a) the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523);

(b) the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523), lacking its associated signal peptide;

(c) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523);

(d) the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523), lacking its associated signal peptide; or

(e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209487.

64. (New) The isolated polypeptide of Claim 63 comprising the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523).

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65. (New) The isolated polypeptide of Claim 63 comprising the amino acid sequence of the polypeptide shown in Figure 222 (SEQ ID NO:523), lacking its associated signal peptide.

66. (New) The isolated polypeptide of Claim 63 comprising the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523).

67. (New) The isolated polypeptide of Claim 63 comprising the amino acid sequence of the extracellular domain of the polypeptide shown in Figure 222 (SEQ ID NO:523), lacking its associated signal peptide.

68. (New) The isolated polypeptide of Claim 63 comprising the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209487.

69. (New) A chimeric polypeptide comprising a polypeptide according to Claim 58 fused to a heterologous polypeptide.

70. (New) The chimeric polypeptide of Claim 69, wherein said heterologous polypeptide is an epitope tag or an Fc region of an immunoglobulin.--

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Applicants respectfully request entry of these new claims for prosecution in this application.  
The Examiner is invited to contact the undersigned at (650) 225-4563 if any issues may be resolved in that manner.

Respectfully submitted,  
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